

1636

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/747,385

DATE: 05/10/2001  
 TIME: 08:37:52

Input Set : A:\Uc-998-1.app  
 Output Set: N:\CRF3\05102001\I747385.raw

#4  
 KFD  
 S-a3-01

ENTERED

3 <110> APPLICANT: Attarian, Gwynne  
 4 Podkaminer, Kara K.  
 5 Yoder, Sean C.  
 6 Kinder Haake, Susan A.  
 7 The Regents of the University of California  
 9 <120> TITLE OF INVENTION: Fusobacterium Nucleic Acids, Plasmids and Vectors  
 11 <130> FILE REFERENCE: 02307E-099810US  
 13 <140> CURRENT APPLICATION NUMBER: US 09/747,385  
 14 <141> CURRENT FILING DATE: 2000-12-22  
 16 <150> PRIOR APPLICATION NUMBER: US 60/173,168  
 17 <151> PRIOR FILING DATE: 1999-12-27  
 19 <160> NUMBER OF SEQ ID NOS: 21  
 21 <170> SOFTWARE: PatentIn Ver. 2.1  
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 24 <211> LENGTH: 407  
 25 <212> TYPE: PRT  
 26 <213> ORGANISM: Fusobacterium nucleatum  
 28 <220> FEATURE:  
 29 <223> OTHER INFORMATION: RepA  
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 36 20 25 30  
 38 His Ile Thr Asn Ile Thr Asn Lys Lys Ile Glu Thr Ile Phe Leu Phe  
 39 35 40 45  
 41 Glu Lys Phe Ile Asn Asp Leu Asp Asn Asn Thr Leu Thr Ile Arg Val  
 42 50 55 60  
 44 Thr Lys Asp Ser Leu Tyr Phe Phe Asn Ile Ala Asn Ser Tyr Leu Arg  
 45 65 70 75 80  
 47 Phe Leu Phe Ser Asp Val Arg Lys Leu Ser Gly Lys Tyr Ser Lys Leu  
 48 85 90 95  
 50 Leu Val Pro Tyr Leu Met Glu Phe Ser His Lys Lys Glu Ala Glu Phe  
 51 100 105 110  
 53 Glu Lys Glu Arg Phe Phe Asn Ile Leu Glu Val Glu Glu Ser Tyr Arg  
 54 115 120 125  
 56 Asn Asn Leu Ser Asp Phe Asn Lys Arg Ile Leu Lys Pro Ala Val Glu  
 57 130 135 140  
 59 Glu Leu Lys Thr Leu Phe Glu Asn Leu Lys Val Glu Arg Leu Lys Asn  
 60 145 150 155 160  
 62 Gly Arg Val Ile Lys Gly Tyr Lys Phe Ser Trp Thr Asn Asp Phe Asn  
 63 165 170 175  
 65 Phe Gln Asn Lys Lys Asp Asn Ile Glu Glu Ala Glu Val Val Glu Glu  
 66 180 185 190  
 68 Lys Glu Asn Ile Ala Ser Gly Glu Leu Glu Lys Tyr Phe Lys Ser Thr  
 69 195 200 205  
 71 Phe Thr Asp Val Asn Tyr Ser Lys Lys His Lys Glu Val Leu Glu Lys

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74 Leu Leu Lys Asn Asn Ser Leu Glu Tyr Ile Lys Lys Tyr Leu Ser Glu
75 225      230      235      240
77 Gln Trp Glu Tyr Val Gln Asn Asp Lys Asn Ile Leu Asn Lys Ser Ala
78      245      250      255
80 Tyr Phe Ser Lys Leu Ile Leu Glu Glu Lys Ala Val Tyr Lys Asn His
81      260      265      270
83 Leu Pro Ala Asp Tyr Glu Glu Leu Lys Val Glu Glu Arg Asn Arg Asn
84      275      280      285
86 Ile Glu Ser Thr Asn Thr Ile Thr Ser Leu Lys Asp Leu Val Glu Lys
87      290      295      300
89 Asp Ile Thr Asp Tyr Glu Val Arg Lys Asn Ile Thr Pro Glu Gln Ile
90 305      310      315      320
92 Glu Gln Glu Val Leu Phe Lys Ile Asp Val Thr Glu Glu Glu Tyr Asn
93      325      330      335
95 Lys Ile Lys Glu Asp Trp Ile Ile Lys Arg Lys Asp Glu Val Pro Asn
96      340      345      350
98 Ser Asp Pro Lys Leu Leu Glu Ile Ile Phe Asn Ala Ser Gln Ser Lys
99      355      360      365
101 Lys Tyr Asn Ile Ile Asn Thr Lys Glu Glu Val Asn Glu Lys Glu Lys
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117 <223> OTHER INFORMATION: repA
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121 tatatttttag gacttcatca aaaattagga aatttacata ttactaatat aacaaataaa 120
122 aaaattgaaa caatcttttt atttgaaaaa ttcataaatg atttagataa taatacttta 180
123 actataagag taacaaaaga ttctctttat tttttaataa ttgctaacag ttatttaagg 240
124 tttctctttt cagatgtagt aaaactttca ggaaaatatt caaagttatt gggttccttat 300
125 ttaatggagt ttagtcataa aaaagaagct gaatttgaaa aagagagatt ttttaattatt 360
126 ctagaagttg aagaaaagta tagaaataat ttatcagatt ttaataagag aattctaaaa 420
127 ccagctgttg aagaattaaa aacacttttt gaaaatttaa aggttgagcg attaaaaaat 480
128 ggaagagtaa taaaaggata taaatttagc tggactaatg attttaattt tcaaaataag 540
129 aaagataata tagaagaagc agaagtagtg gaagaaaaag aaaatattgc ttcaggagag 600
130 ttagaaaaat attttaaatc aacttttact gatgtaaatt attcaaagaa gcataaagaa 660
131 gtttttagaaa aattattaaa aaataatagt ttagaatata ttaaaaaata tttatctgag 720
132 cagtgggagt atgtacaaaa tgataaaaaa attttaaata aatcagcata tttctcaaaa 780
133 ctaatttttag aagaaaaagc agtatataaa aatcatctac cagctgacta tgaagaacta 840
134 aaagttgaag aaagaaatag aaatatagaa agtacaaata ctattacatc attaaaagat 900
135 ttagtagaaa aagacattac agattatgaa gttagaaaga atataactcc tgaacaaata 960
136 gaacaagaag ttttatttaa aatagatgta actgaagaag aatataataa gattaaagaa 1020

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137 gattggataa taaaacgaaa agatgaagtt cctaatagtg atccaaaact tttagaaatt 1080
138 atattttaatg caagtcaatc aaaaaaatat aatataatta atactaaaga agaagttaat 1140
139 gaaaaagaaa aagagcttca cgaattagaa gaaaatataa aaagaatgca agaagaacta 1200
140 aataaattaa aaaaagaggt atag 1224
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144 <211> LENGTH: 22
145 <212> TYPE: DNA
146 <213> ORGANISM: Artificial Sequence
148 <220> FEATURE:
149 <223> OTHER INFORMATION: Description of Artificial Sequence: iteron sequence
150 within the origin of replication of plasmid pFN1
152 <400> SEQUENCE: 3
153 tcaactttaa caggacaaat tt 22
156 <210> SEQ ID NO: 4
157 <211> LENGTH: 132
158 <212> TYPE: DNA
159 <213> ORGANISM: Artificial Sequence
161 <220> FEATURE:
162 <223> OTHER INFORMATION: Description of Artificial Sequence: six copies of
163 the iteron sequence within the origin of
164 replication of plasmid pFN1
166 <400> SEQUENCE: 4
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168 aaattttcaa cttaacagg acaattttc aactttaaca ggacaaattt tcaactttaa 120
169 caggacaaat tt 132
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173 <211> LENGTH: 1230
174 <212> TYPE: DNA
175 <213> ORGANISM: Artificial Sequence
177 <220> FEATURE:
178 <223> OTHER INFORMATION: Description of Artificial Sequence: repA homolog
179 sequence of plasmid pAD52
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184 aaaattgaaa caatcttttt atttgaaaaa ttcataaatg atttagataa taatacttta 180
185 actataagag taacaaaaga ttctctttat ttttttaata ttgctaacag ttatttaagg 240
186 tttctctttt cagatgtagt aaaactttca ggaaaatatt caaagttatt gggttccttat 300
187 ttaatggagt ttagtcataa aaaagaagct gaatttgaaa aagagagatt ttttaatat 360
188 ctagaagttg aagaaaagta tagaataaat ttatcagatt ttaataagag aattctaaaa 420
189 ccagctgttg aagaattaaa aacacttttt gaaaatttaa aggttgagcg attaaaaaat 480
190 ggaagagtaa taaaaggata taaatttagc tggactaatg attttaattt tcaaaataag 540
191 aaagataata tagaagaagc agaagtagtg gaagaaaaag aaaataaaaa tattgctcct 600
192 ggagagttag aaaaatattt taaaacaact ttccctggtg taaattattc aaagaagcat 660
193 aaagaagttt tagaaaaatt attaaaaaat aatagtttag aatatattaa aaatatttta 720
194 tctgagcagt gggagtatgt acaaaacgat aaaaatattt taaataaatc agcatatttt 780
195 tcaaaactaa tcttagaaga aaaagcagta tataaaaatc atctaccagc tgactatgaa 840
196 gaattaaaag ttgaagaaag aaatagaaat atagaaagta caaatactat tacatcatta 900
197 aaagatttag tagaaaaaga cattacagat tatgaattta gaaagaatat aactcctgaa 960

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198 caaatagaac aagaagtttt atttaaaata gatgtaactg aagaagaata taataagatt 1020
199 aaagaagatt ggataataaa acaaaaagaa gtagttccta atagtgatcc agaactttta 1080
200 gaagttatat ttaatgcaag tcaatcaaaa aaatataata taattaatac taaagaagaa 1140
201 gttaatgaaa aagaaaaaga gcttcacgaa ttagaagaaa atataaaaag aatgcaagaa 1200
202 gaaataaata aattaaaaaa agaggtatag                                     1230
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206 <211> LENGTH: 5887
207 <212> TYPE: DNA
208 <213> ORGANISM: Artificial Sequence
210 <220> FEATURE:
211 <223> OTHER INFORMATION: Description of Artificial Sequence: plasmid pFN1
213 <400> SEQUENCE: 6
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215 tttttactttt ttcaattttt tctaattcaa tcgcttttaa ttctttctaaa gtcttttttaa 120
216 ttttttttagt agcttccata cataatcacac tccagcatta ttatttataa aaatataatt 180
217 atataataca tatctagttaa aataaatcaa gtagtgcggg cttaaacaaag agccatataa 240
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222 tgcaattatt ttataaaacc aactttaaag agagtataaa taatgattaa atttacatta 540
223 agattaataa taatgattaa atttacatta agattaacgg aagatgaaaa aaaactttta 600
224 gatataaaag ctgatgaatt aggtaaatca aaaaatgaag ttttaaagtt tottataaac 660
225 aataaattgg aagatactaa aaaagaattt gacctattaa atgagcttga taaaaattat 720
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239 attttataaa agcattagat gaaaaagggtg ttattgtgga ttgggaagac cataaaaaac 1560
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246 aaatgtgaat gagcaagaaa taaaaatgga tagttatatt aaaaaataa ttgaaaatgt 1980
247 tttagaagtt cagctaaaag aacataaaga aatagcttcc attgctaaaa ctaaaatagc 2040
248 tgaagtaact ttagaactag aaaaattaaa acagctggag aaagcaacta ctaaaatagc 2100
249 agatgataca aatataatta caaataaaat gattgaaaat gttgaaaatt ataataaagt 2160
250 atttttagaa agaattgata aatttaattt attgatggta gaaaagttaa atgaagtaaa 2220

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253 tacttttaaat ttaattgaag atagtctaaa taaattaaaa atagccttct atacagctgt 2400
254 atctgttatt gtaatatattt tatttttttac ggggataatt ttatataaga caaataatag 2460
255 agttgctagt gttgaagaaa gcttaataaa tatatctagt tcagtaactg gattagttaa 2520
256 aggggactta aagttttggt acagtgaaga agacaaaaaa gcctatgtaa gtaatgtaga 2580
257 aagtattaaa aaaaaataag atagcaagaa gaaaaataa aagcttcagt aagataaaaa 2640
258 agcaataaac atttaattta ttgctttttt attttatagt ttagtcattg agggtaaatt 2700
259 tttatagtaa tatatatattt aacaatttta ctatattact ttttaacatt ctttagaaac 2760
260 atatccataa tatagttcat tagacttgcg acagttatct catttgtagc agcatacttt 2820
261 ttgaaatttg agtaaattctc tgagtttggt ttcatcgata tagttatatc atttttttaa 2880
262 gttctataat gttcaggaa gacaacagtt tcattattta cattaatttt tctttcgtta 2940
263 acaatattaa aaagtatttc taaattttct tcattaaaca aactattata tttttctgga 3000
264 actgtcaatt gaaattcttg cttttcattt atattacttt ttatattact atctatatta 3060
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278 gcagttatat taatatgggt tcacattatc acaagtttat tagcatagtt cttgtaaaaa 3900
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299 aggagagtta gaaaaattt ttaaatcaac ttttactgat gtaaattatt caaagaagca 5160

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**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

## VERIFICATION SUMMARY

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Input Set : A:\Uc-998-1.app

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L:428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:432 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:433 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:436 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14